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Scoring table:
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    Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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Match Length DB
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Listing first 45 summaries
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/cgn2_6/ptodata/1/laa/5B_COMB.pep:*
/cgn2_6/ptodata/1/laa/6A_COMB.pep:*
/cgn2_6/ptodata/1/laa/6B_COMB.pep:*
/cgn2_6/ptodata/1/laa/B_COMB.pep:*
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US-09-268-5448-36
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US-09-138-277C-3
US-09-138-277C-3
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US-08-744-139-4
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US-08-691-8148-2
US-08-691-8148-29
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4.4	4.4	4.4	4.4	4.4	4.6	4.7	4.8	4.8	4.8	4.8	4.8		6.2		8.3	8.6	9.8
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sequence I/, Appr	Sequence 17 Appr	Sequence 1/, Appr	Sequence 10, Appr	Sequence 4, Appril	· _	. 0	Ļ	· -	٠ د	Sequence 5, Appri	7) (A	,	2 0	2 5	3 5	Sequence 30, Appl

ALIGNMENTS

US-08-697-610-2 Sequence 2, Application US/08697610; Patent No. 6172187 ; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-697-610-2 GENERAL INFORMATION: PRIOR APPLICATION DATA: APPLICATION NUMBER: US 08/349,357 APPLICATION NUMBER: US 08/349,357 FILLING DATE: 02-DEC-1994 ATTORNEY/AGENT INFORMATION: NAME: Campbell, Cathryn A. REGISTRATION NUMBER: 31,815 REGISTRATION NUMBER: P-LJ 1203 TELECOMMUNICATION INFORMATION: TELEPHONE: (619) 535-9001 TELEPAX: (619) 535-9001 INFORMATION FOR SEQ ID NO: COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS TITLE OF INVENTION: CD40 Associated Proteins NUMBER OF SEQUENCES: 17 APPLICANT: Reed, John C. APPLICANT: Sato, Takaaki SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/697,610 CORRESPONDENCE ADDRESS: SEQUENCE CHARACTERISTICS: LENGTH: 543 amino acids COUNTRY: STREET: ADDRESSEE: CLASSIFICATION: FILING DATE: 92122 San Diego amino acid California E: Campbell and Flores 4370 La Jolla Village I USA 530 Village Drive, Suite 700 P-LJ 1203

541;

Conservative

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Mismatches

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Indels Length 543;

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Gaps

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94.1%; 95.2%;

Score 2831.5; DB 4; Pred. No. 2.2e-245;

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                               TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
SEQUENCE CHARACTERISTICS:
LENGTH: 543 amino acids
                                                                    REFERENCE/DOCKET NUMBER: P-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,81
                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/349,357
                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Sato, Takaaki
TITLE OF INVENTION: CD40 Associated Proteins
NUMBER OF SEQUENCES: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Reed, John C. APPLICANT: Sato, Takaaki
                                                                                                                                                                FILING DATE: 02-DEC-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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STATE: Californi
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                                                                                                                                                                                                                                                                                                                                                                        California
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                                                   : (619) 535-9001
(619) 535-8949
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                                                                                                                      31,815
                                                                                                       P-LJ 1203
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                                                                                                                                            SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                       Query Match
     Matches 438;
                    Best Local Similarity
                                                                                                                                                                                                                                                                                                                           Sequence 36, Application US/09268544B Patent No. 6410710
                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/268,544B
CURRENT FILING DATE: 1999-03-11
NUMBER OF SEQ ID NOS: 43
                                                                                                                                                                                                                                APPLICANT: Lederman, Seth
APPLICANT: van Eyndhoven, Winfried
TITLE OF INVENTION: TRAF-3 Deletion Isoforms and Uses Thereof
FILE REFERENCE: 0575-58732
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                                                                                                              TYPE: PRT
                                                                                           ORGANISM: Human
                                                                                                                             LENGTH: 438
                                                                                                                                                                                                                                                                                                                                                                                                                                516 TVLENGTYIKDDTIFIKVIVDTSDLPDP 543
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              74.8%; Score 2251; DB 4; Length 438; 77.1%; Pred. No. 1.7e-193;
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95.2%; Pred. No. 2.2e-245;
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APPLICANT: van Eyndhoven, Winfried
APPLICANT: van Eyndhoven, Winfried
TITLE OF INVENTION: TRAF-3 Deletion Isoforms And Uses Thereof
FILE REFERENCE: 0575-58732
CURRENT APPLICATION NUMBER: U5/09/268,544B
CURRENT FILING DATE: 1999-03-11
NUMBER OF SEQ ID NOS: 43
SOFTWARE: Patentin Ver. 2.1
SOFTWARE: Patentin Ver. 2.1
LENGTH: 347
THEREFORE SEQ ID NOS: 43
LENGTH: 347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 38, Application US/09268544B Patent No. 6410710 GENERAL INFORMATION:
                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                              Matches 346; Conservative
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121 SRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHC 180
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                                  CSPKQTECGHRECESCMAALLSSSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNE 120
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                                                                                                       MESSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKTVEDKYKCEKCHLVL 60
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                                                                                                                                                                       56.3%; Score 1692.5; DB 4; Length 347; 60.9%; Pred. No. 1.4e-143; ative 0; Mismatches 1; Indels 221;
                                                                                                                                                                         1; Indels 221; Gaps
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180	130 LGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKYREATCSHCKSQVPMIAL 1	Оy	
121	71 RFCESCMAALLS-SSSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNESRGCAEQLT 1	ОУ	
70 62	22 HTDRSAGTP-VFVPEQGGYKEKFVKTVEDKYKCEKCHLVLCSPKQTECGH 7 : : : :	Db	
os 10;	uery Match 41.7%; Score 1253.5; DB 4; Length 558; sest Local Similarity 41.6%; Pred. No. 6.2e-104; satches 241; Conservative 114; Mismatches 166; Indels 59; Gaps	Que Bes Mat	
7 199 300 360 139 199 480 540 319	FOGTNOQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI FOGTNOQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI FOGTNOQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI FOGTNOQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEI FORTNOQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLKEIDKEITPFRQNWEEADSWKSSVESLQNR FORTNOCH HILL	טוואס מיו מסימו	
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APPLICANT: NAKATA, MOTOMI

APPLICANT: NAKANO, HIROYASU

APPLICANT: NAKANO, HIROYASU

APPLICANT: OKUMURA, KO

TITLE OF INVENTION: TRAF FRAMILY MOLECULES, POLYNUCLEOTIDES ENC.

TITLE OF INVENTION: AND ANTIBODIES AGAINST THEM

FILE REFERENCE: 007898-0255515

CURRENT APPLICATION NUMBER: US/09/138,277C

CURRENT FILING DATE: 1998-08-18

PRIOR APPLICATION NUMBER: POT/JP97/00512

PRIOR FILING DATE: 1997-02-24

PRIOR APPLICATION NUMBER: JP 34674/1996

PRIOR APPLICATION NUMBER: JP 34674/1996

PRIOR FILING DATE: 1996-02-22

NUMBER OF SED ID NOS: 16

SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 3
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204 CPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCVFQGTNQQIKAHEASSAVQHVNLL 263
                                           135 QPVQCSNEKCREPVLRKDLKEHLSASCQFRKEKCLYCKKDVVVINLQNHEENLCPEYPVF 194
                                                                 Local Similarity
hes 232; Conserv
                                                                                                                                                                                                                     17 QNSGNSISLDFEPSIEYQFVERLEERYKCAFCHSVLHNPHQTGCGHRECQHCILSLRELN 76
                                                                                                                                                                                                                                         LVDAVDSVKQRITQLEASD------QRLVLLEGETSKHDAHINIHKAQLNKNEERFKQL
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                                                                                                                              TVPICPVDKEVIKSQEVFKDNCCKREVLNLYVYCSN-APGCNAKVILGRYQDHLQ-QCLF 134
                                                                                                                                                                          SSPKCTACQESIVKDKVFKDNCCKREILALQIYCRNESRGCAEQLTLGHLLVHLKNDCHF 143
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                                                                                                                                                                                                                                                                                                           Conservative 121;
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                                                                                                                                                                                                                                                                                                                           39.6%; Score 1192.5; DB 4; Length 557; 42.0%; Pred. No. 1.8e-98;
                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ENCODING
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RESULT 8 US-08-331-394-4

Sequence

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                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 228
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/09181958 Patent No. 6143507
                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Kehry, Marilyn R
APPLICANT: Pullen, Steven S
APPLICANT: Crute, James J
TITLE OF INVENTION: High Throughput Compatible Assays for Receptor-TRAF
TITLE OF INVENTION: Interactions
FILE REFERENCE: 9_142_Nucl.Seq
CURRENT APPLICATION NUMBER: US/09/181,958
CURRENT FILING DATE: 1998-10-29
NUMBER OF SEQ ID NOS: 2
                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: human
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                  534 CPVFVAQTVLENGTYIKDDTIFIKVIVDTSDLPDP 568
                                                                                                                                                                                                                                 354 VESLQNRVTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETAS 413
181
                                                                      121
                                                                                                                                          61 YNGYLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSL
                                                                                                                                                                                                                                                                                  Local Similarity
wes 215; Conserv
                                                                                                                                                                                                                 1 VESLQNRVTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LEKNVQLEEQISDLHKSLEQKESKIQQLAETIKKLEKEFKQFAQLFGKNGSFLPNIQ-VF
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                                                                                       FFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASG
                                                                                                                                                            YNGVLIWKIRDYKRRKOEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSL 473
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CPVFVAQTVLENGTYIKDDTIFIKVIVDTSDLPDP
                                                                   FFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASG
                                                                                                                                                                                                                                                                                  37.2%; Score 1118; DB 4; ilarity 100.0%; Pred. No. 2.3e-92; Conservative 0; Mismatches 0:
                                                                                                                                                                                                                                                                                                                         Length 228;
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APPLICANT: Rothe, Mike
TITLE OF INVENTION: Tumor Necrosis Factor
TITLE OF INVENTION: Receptor-Associated Factors
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
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GENERAL INFORMATION:
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NAME: Dreger, Ginger ...

REGISTRATION NUMBER: 33,055

REFERENCE/DOCKET NUMBER: 897P

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 1.5,725-2316
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/250
FILING DATE: 27-MAY-1994
ATTORNEY/AGENT INFORMATION:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/3:
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MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 501 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                               113 LQIYCRNESRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKY 172
                                                                                                                                                                                                                                   233 SFKRYGCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVS------LLQNESV 282
                                                                                                                                                                                                                                                                                                                                                                                  103 LPAVCPND--GCTWKGTLKEYESCHEGLCPFLLTEC--PACKGLVRLSEKEHHTEQECPK 158
                                        343 NWEEADSMKSSVESLONRVTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADM 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 415/225-32:
TELEFAX: 415/952-9881
TELEX: 910/371-7168
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2 AAASVTSPGSLELLQP-------GFSKTLLGTRLEAKYLCSACKNILR 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3 SSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKT-VEDKYKCEKCHLVLC 61
                                                                                                                                         EKNKSIQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQ 342
                                                                                                                                                                                                                                                                                                                                   REATCSHCKSQVPMIALOKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RPFQAQCGHRYCSFCLTSILSSGPQNCAACVYEGLYEEGISILESSSAFPDNAARREVES 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPKQTECGHRFCESCMAALLSSSSPKCTAC-----QESI---VKDKVFKDNCCKREILA 112
-----KIEALSNKVQQLE-----RSIGL------KDLAMADL
                                                                                               LEQK-IATFENIVCVLNREVER-
                                                                                                                                                                                         REHTVGCSEMVETENLODHELQRLREHLALL -- LSSFLEAQASPGTLNQVGPELLQRCQI 274
                                                                                                                                                                                                                                                                                     RSLSCQHCRAPCSHVDLEVHYEV-CPKFPLTC-DGCGKKKIPRETFQDHVRACSKCRVLC
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                                                                                                 --VAVTAEACSRQHRLDQD-----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: 5.25 inch, 360 Kb:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 501 amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: 89
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors NUMBER OF SEQUENCES: 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goeddel, David V. APPLICANT: Rothe, Mike
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     463 DGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFK 522
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173 REATCSHCKSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                  103 LPAYCPND--GCTWKGTLKEYESCHEGLCPFLLTEC--PACKGLVRLSEKEHHTEQECPK 158
                                                                      113 LQIYCRNESRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKY 172
                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 415/--
TELEFAX: 910/371-7168
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CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
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                                                                                                              43 RPFQAQCGHRYCSFCLTSILSSGPQNCAACVYEGLYEEGISILESSSAFPDNAARREVES 102
                                                                                                                                                    62 SPKQTECGHRFCESCMAALLSSSSPKCTAC----QESI---VKDKVFKDNCCKREILA 112
                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                          3 SSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKT-VEDKYKCEKCHLVLC 61
                                                                                                                                                                                          2 AAASVTSPGSLELLQP-------GFSKTLLGTRLEAKYLCSACKNILR 42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RPVSDMNIASGCPLFCPVSKMEAKNSYVRDDAIFIKAIVDLTGL 501
                                                                                                                                                                                                                                                                    196;
                                                                                                                                                                                                                                                                  similarity 33.6
96; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  415/225-3216
                                                                                                                                                                                                                                                                                      27.8%; Score 837.5; DE 33.6%; Pred. No. 1e-66;
                                                                                                                                                                                                                                                                      ; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        360 Kb floppy disk
                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                        194; Indels 105;
                                                                                                                                                                                                                                                                                                        Length 501;
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US-08-446-915-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4,
                                                                 TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 K
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-D
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                           TELEPHONE: 415/225-3216
                                                                                                                                                                                                                                    FILING DATE: 28-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Goeddel, David V. APPLICANT: Rothe. Mike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 460 Point San Bruno Blvd CITY: South San Francisco
STRIE: California
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                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             463 DGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 339 EQKVSELEVSTYDGVFIWKISDFTRKRQEAVAGRTPAIFSPAFYTSRYGYKMCLRVYLNG
               TOPOLOGY:
                                                                                                                          TELEFAX:
                                                                                                                                                                            REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                 NAME: Dreger, Ginger REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/2: FILING DATE: 27-MAY-1994
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                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SFKRYGCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVS------LLQNESV 282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RSLSCQHCRAPCSHVDLEVHYEV-CPKFPLTC-DGCGKKKIPRETFQDHVRACSKCRVLC 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94080
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 196;
                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44 mb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: TUMOR NECROSIS NUMBER OF SEQUENCES: 59
                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Goeddel, David V APPLICANT: Rothe, Mike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          399
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                                                 APPLICATION NUMBER: US/0 FILING DATE: 31-Oct-1996
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 APPLICATION NUMBER:
                                                                                                                                                                                                    ZIP:
                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                        STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              113 LQIYCRNESRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKY 172
                                                                                                                                                                                                                                                                                         ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DGMGKGTHLSLFEVIMRGEYDALLPWPFKQKYTLMLMDQGSSRRHLGDAFKPDPNSSSFK 522
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460 Point San Bruno Blvd
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                                                                     US/08/744,139
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                                                                                                                                                                                                                           RESULT 12
PCT-US95-06639-4
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                                                                                                                                                                               Sequence 4, Application PC/TUS9506639 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 501 amino acids
TYPE: Amino Acid
  CORRESPONDENCE ADDRESS:
ADDRESSE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                        TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: 05/27/1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                     458
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     173
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      43
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REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: P0897C1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62 SPKQTECGHRFCESCMAALLSSSSPKCTAC-----QESI---VKDKVFKDNCCKREILA 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 27.8%; Score 837.5; DE Local Similarity 33.6%; Pred. No. 1e-66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N
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                                                                                                                                                                                                                                                                                                                                     KPTGEMNIASGCPVFVAQTVLE-NGTYIKDDTIFIKVIVDTSDL 565
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DLRFQVLETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNG 462
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NWEEADSMKSSVESIQNRVTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADM 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REATCSHCKSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                                                                                                                                                                                                                                                                                                   RPVSDMNIASGCPLFCPVSKMEAKNSYVRDDAIFIKAIVDLTGL 501
                                                                                                                                                                                                                                                                                                                                                                                                                                                DGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFK 522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EQKVSELEVSTYDGVFIWKISDFTRKRQEAVAGRTPAIFSPAFYTSRYGYKMCLRVYLNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -----KIEALSNKVQQLE------RSIGL-----KDLAMADL 338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LEQK-IATFENIVCVLNREVER----- 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EKNKSIQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQ 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SFKRYGCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVS------LLQNESV 282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RSLSCOHCRAPCSHVDLEVHYEV-CPKFPLTC-DGCGKKKIPRETFQDHVRACSKCRVLC 216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LQIYCRNESRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKY 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RPFQAQCGHRYCSFCLTSILSSGPQNCAACVYEGLYEEGISILESSSAFPDNAARREVES 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAASVTSPGSLELLQP-------GFSKTLLGTRLEAKYLCSACKNILR 42
                                                                                                                                                                                                                                                                                                                                                                                                      DGTGRGTHLSLFFVVMKGPNDALLQWPFNQKVTLMLLDH-NNREHVIDAFRPDVTSSSFQ 457
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LPAVCPND--GCTWKGTLKEYESCHEGLCPFLLTEC--PACKGLVRLSEKEHHTEQECPK 158
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 501 amino acid
TYPE: amino acid
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/25-3216
TELEPAX: 415/952-9881
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FILING DATE: 28-OCT-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Ginger R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch, 1.44
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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CURRENT APPLICATION DATA:
                                                                                                                                                        403 DLRFQVLETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNG
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523 KPTGEMNIASGCPVFVAQTVLE-NGTYIKDDTIFIKVIVDTSDL 565
                                                                                                                                                                                                                                                                  343 NWEEADSMKSSVESLQNRVTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADM 402
                                                                                                                                                                                                                                                                                                                                                                                                  217 RFHTVGCSEMVETENLQDHELQRLREHLALL--LSSFLEAQASPGTLNQVGPELLQRCQI 274
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               103 LPAVCPND-:GCTWKGTLKEYESCHEGLCPFLLTEC--PACKGLVRLSEKEHHTEQECPK 158
                                                                                                                                                                                                                                                                                                                                                      283 EKNKSIQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQ 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       43 RPFQAQCGHRYCSFCLTSILSSGPQNCAACVYEGLYEEGISILESSSAFPDNAARREVES 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62 SPKQTECGHRFCESCMAALLSSSSPKCTAC-----QESI---VKDKVFKDNCCKREILA 112
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REGISTRATION NUMBER: 33,055
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: PCT/US95/06639
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 AAASVTSPGSLELLQP-------GFSKTLLGTRLEAKYLCSACKNILR 42
                                         DGTGRGTHLSLFFVVMKGPNDALLQWPFNQKVTLMLLDH-NNREHVIDAFRPDVTSSSFQ
                                                                              DGMGKGTHLSLEFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFK 522
                                                                                                                                   EQKVSELEVSTYDGVFIWKISDFTRKRQEAVAGRTPAIFSPAFYTSRYGYKMCLRVYLNG
                                                                                                                                                                                                                                                                                                            LEQK-IATFENIVCVLNREVER------VAVTAEACSRQHRLDQD-----
                                                                                                                                                                                                                                                                                                                                                                                                                                             SFKRYGCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVS------LLQNESV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REATCSHCKSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LQIYCRNESRGCAEQLTLGHLLVHLKNDCHFEELPCVRPDCKEKVLRKDLRDHVEKACKY 172
                                                                                                                                                                                                                      -----KIEALSNKVQQLE-----RSIGL------KDLAMADL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27.8%; Score 837.5; DB 5; 33.6%; Pred. No. 1e-66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inch, 1.44 Mb disk
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458 RPVSDMNIASGCPLFCPVSKMEAKNSYVRDDAIFIKAIVDLTGL

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 409 amino acic
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SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: Tumor Necrosis Factor TITLE OF INVENTION: Receptor-Associated Factors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 89
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 27-MAY-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Goeddel, Day APPLICANT: Rothe, Mike
  445 FYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSS 504
                                         229 VVELQQTLAQKDQVLGKLEHSLRLMEEASFDGTFLWKITNVTKRCHESVCGRTVSLFSPA
                                                                               389 ----DQMLSVHDIRLADMDLRFQVLETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQP 444
                                                                                                                                                                    337 IRPFRQNWEEADSMKSSVESLQNRVTELESVDKSAGQVARNTGLLESQLSRH-----
                                                                                                                                                                                                                                                        278 QNESVEKNKSIQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEK-LKELDKE 336
                                                                                                                                                                                                                                                                                                                                           233 SFKRYGCVFQGTNQQIKAHEASSAVQHVNL----LKEWSNS-------LEKKVSLL 277
                                                                                                                                                                                                                                                                                                                                                                                                                               173 REATCSHCKSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                                                                                                                                                                                                                                                                                                                                                                     28 RVLCCTACLSE----NLRDDEDRICPKCRADNLHPVSPGSPLTQE-KVH-SDVAEAEIMC 81
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: South San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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                                                                                                                                                                                                              Q-----LQAAVEATGDLEVDCYRAPCCESQEELALQHLVK-----EKLLAQLEEK 186
                                                                                                                                                                                                                                                                                                    PFAGVGCSFKGSPQSMQEHEATSQSSHLYLLLAVLKEWKSSPGSNLGSAPMALERNLSEL 141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       154;
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460 Point San Bruno Blvd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                             -----ANIVAVLNKEVE-----ASHLALAASIHQSQLDREHLLSLEQR 228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08/250858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         69; Mismatches 128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         897P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 628; DB 1;
Pred. No. 4.5e-48;
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US-08-250-858-2
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                                                                                                                                                                                                                                                                Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2,
                                                                                                                                                                                                                                                Matches 154;
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                                                                                                                                                                                                                                                                                                                                                                                                      TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-DOS/MS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,055
REFERENCE/DOCKET NUMBER: 89
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
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337 IRPFRONWEEADSMKSSVESLONRVTELESVDKSAGOVARNTGLLESOLSRH------
                                  142
                                                         278 QNESVEKNKSIQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEK-LKELDKE 336
                                                                                                                                       233 SFKRYGCVFQGTNQQIKAHEASSAVQHVNL----LKEWSNS-------LEKKVSLL 277
                                                                                                                                                                                                            173 REATCSHCKSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                                                                                                                                                                                                                                                                                                                   TYPE: ami
TOPOLOGY:
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                                                                                                       82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 460 Point San Bruno Blvd CITY: South San Francisco
                                                                                                                                                                          28 RVLCCTACLSE----NLRDDEDRICPKCRADNLHPVSPGSPLTQE-KVH-SDVAEAEIMC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
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                                                                                                                                                                                                                                                                                                                                                      amino acid
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                                                                                                                                                                                                                                                                                                                                                                      409 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                            415/952-9881
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                                                                                                                                                                                                                                                Conservative
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SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                     linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-May-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumor Necrosis Factor Receptor-Associated Factors
                                                                                                                                                                                                                                              20.9%; score 628; DB 1; 1 36.6%; Pred. No. 4.5e-48; tive 69; Mismatches 128;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              David V.
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US-08-446-915-2
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                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/250858
FILING DATE: 27-MAY-1994
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNMBER: 08/331394
FILING DATE: 28-0CT-1994
ATTORNEY/AGENT INFORMATION:
NAME: DIEGET, Gliger R.
REGISTRATION NUMBER: 897P:
REFERENCE/DOCKET NUMBER: 897P:
TELECOMMUNICATION INFORMATION:
TELEPAX: 415/252-3316
TELEFAX: 415/252-9881
INFORMATION GY SEG ID NO: 2:
REGULENCE CLARACTERITICS: 2:
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                                                                                                           Query Match
Best Local :
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                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 409 amino acids
TYPE: amino acid
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MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,915
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Goeddel, David v.
APPLICANT: Rothe, Mike
TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors
NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
                                   173 REATCSHCKSQVPMIALQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTC 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: CE
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28 RVLCCTACLSE----NLRDDEDRICPKCRADNLHPVSPGSPLTQE-KVH-SDVAEAEIMC 81
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STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
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                                                                                                         Local
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                                                                                  Similarity 36.6
54; Conservative
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                                                                                                                                                                                       linear
                                                                             20.9%; Score 628; DB 1; Length 409; 36.6%; Pred. NO. 4.5e-48; ative 69; Mismatches 128; Indels
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408 S 408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    278
                                                                                                                                                                                                                                         229 VVELQQTLAQKDQVLGKLEHSLRLMEEASFDGTFLWKITNVTKRCHESVCGRTVSLFSPA 288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      82 PFAGVGCSFKGSPQSMQEHEATSQSSHLYLLLAVLKEWKSSPGSNLGSAPMALERNLSEL
                                                                                                                                                                               FYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSS 504
                                                                          REHAIDAFRPDLSSASFQRPQSETNVASGCPLFFPLSKLQSPKHAYVKDDTMFLKCIVDT 407
                                                                                                                                                          FYTAKYGYKLCLRLYLNGDGSGKKTHLSLFIVIMRGEYDALLPWPFRNKVTFMLLDQ-NN
                                                                                                                                                                                                                                                                                                                       LRVF-----ANIVAVLNKEVE-----ASHLALAASIHQSQLDREHLLSLEQR 228
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Search completed: December 19, 2002, 14:56:21 Job time: 18 secs

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